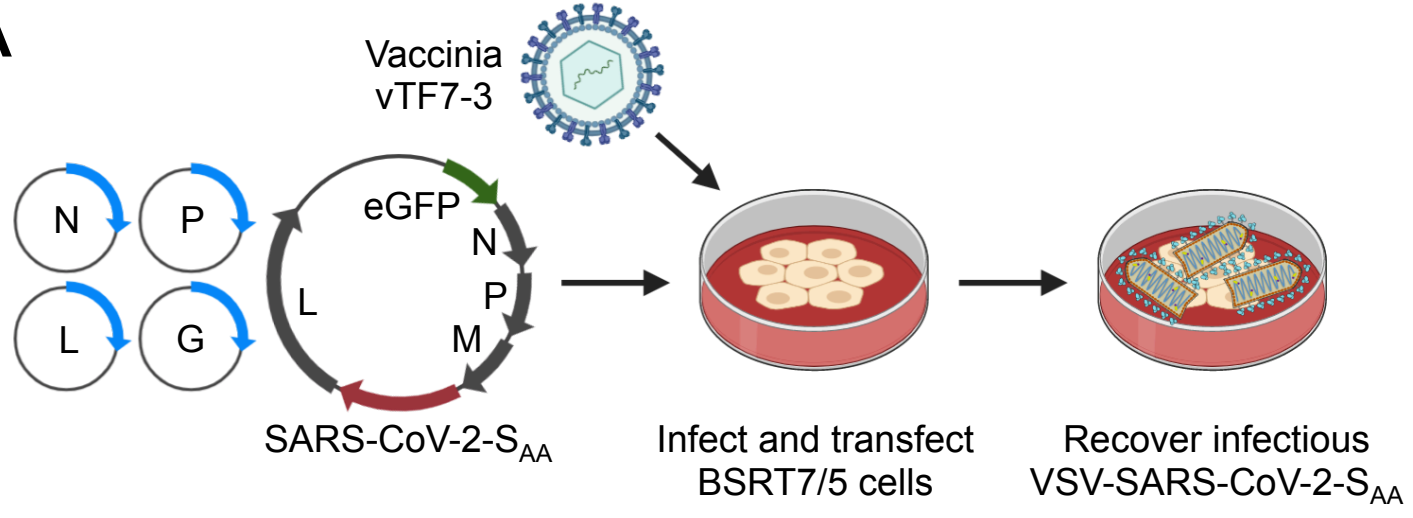


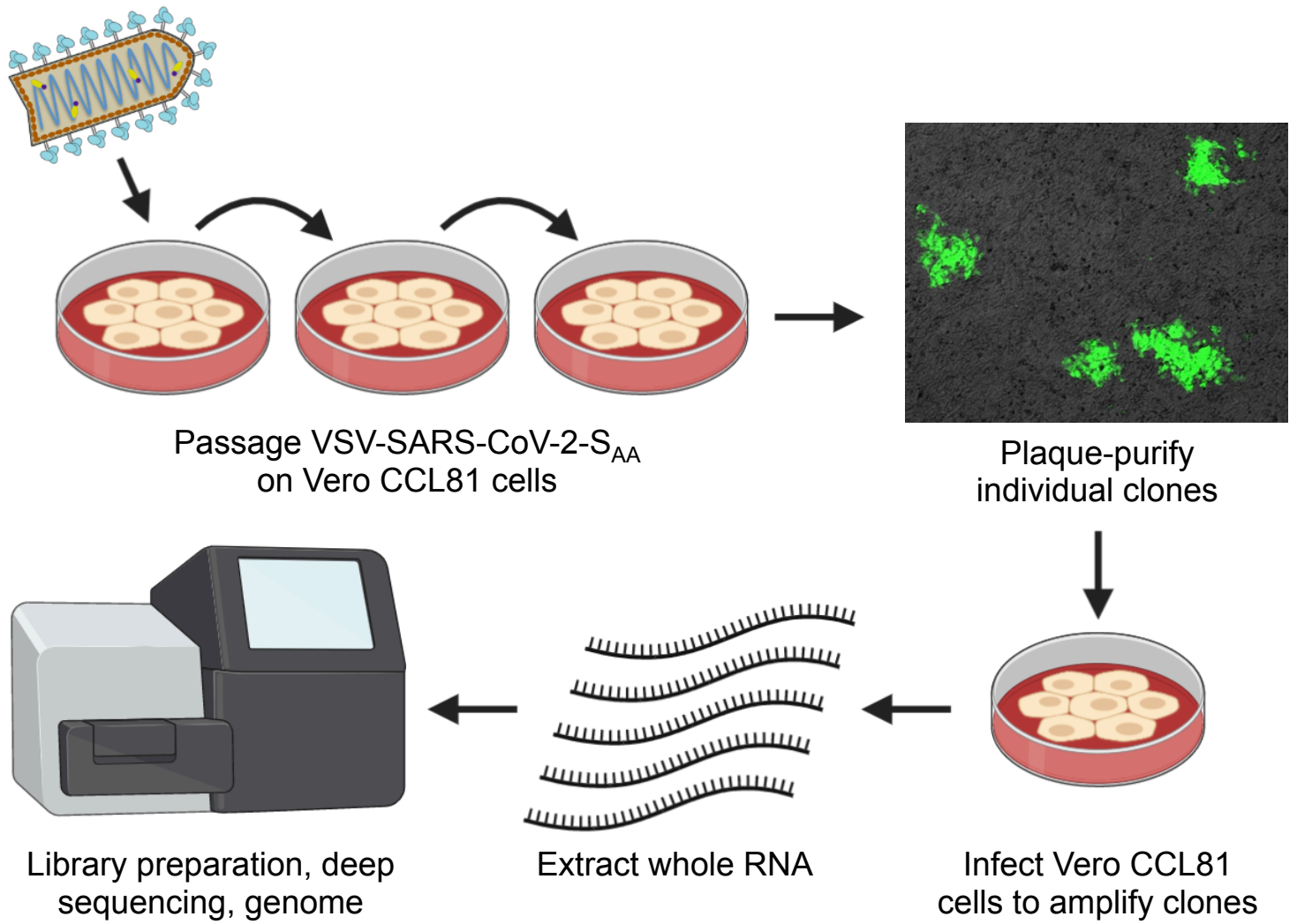
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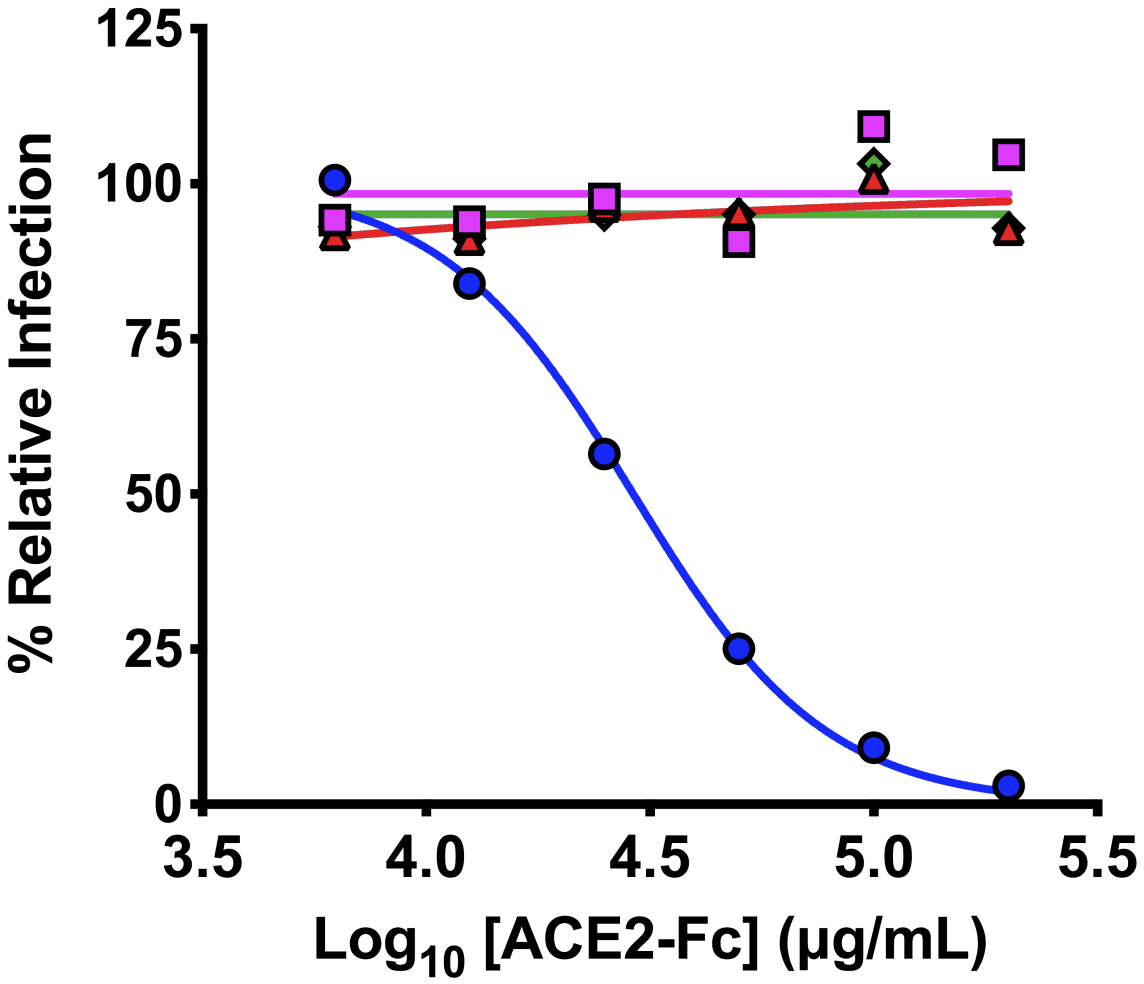


B

Tail Mutant	Amino acid sequence (membrane proximal region, transmembrane domain, cytoplasmic tail)	Rescue	Spread
S _{AA}	LNESLIDLQELGKYEQYIKWPWYIWLGFIAGLIAIVMVTIMLCCMTSCCCLKGCCSCGSCCKFDEDDSEPVLKGVALAYT	+	+
MERS S _{AA}	LNESLIDLQELGKYEQYIKWPWYIWLGFIAGLIAIVMVTIMLKLKCNRCDDRYEEDLEPHAVAVH	+	-
VSV G #1	LNESLIDLQELGKYEQYIKWPWYIWLGFIAGLIAIVMVTIMLCCMTSCCCLRVGIYLCIKLKHTKKRQIYTDIEMNRLGK	+	-
VSV G #2	LNESLIDLQELGKYEQYIKWPWYIWLGFIAGLIAIVMVTIMLRVGIYLCIKLKHTKKRQIYTDIEMNRLGK	+	-
VSV G TM/tail	LNESLIDLQELGKYEQYIKWSSIASFCFIIGLIIGLFLVLRVGIYLCIKLKHTKKRQIYTDIEMNRLGK	+	-
VSV G Ecto/TM/tail	LNEGWFSSWKSSIASFCFIIGLIIGLFLVLRVGIYLCIKLKHTKKRQIYTDIEMNRLGK	+	-

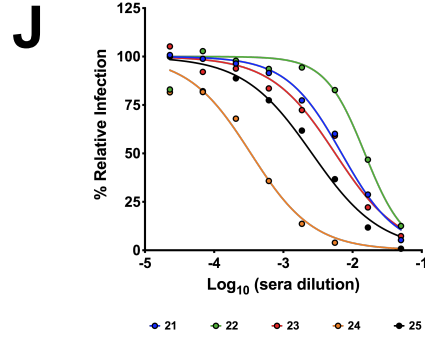
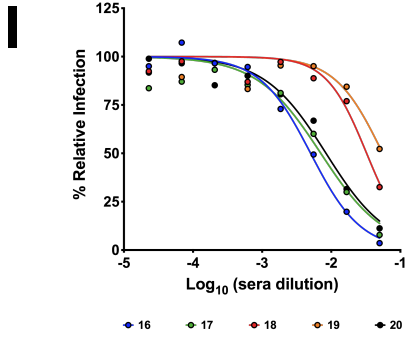
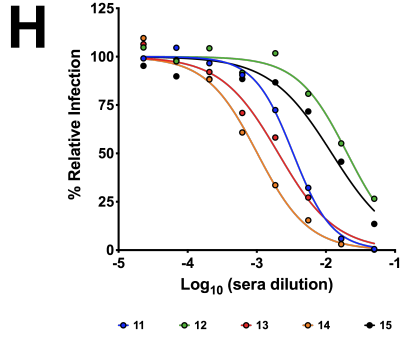
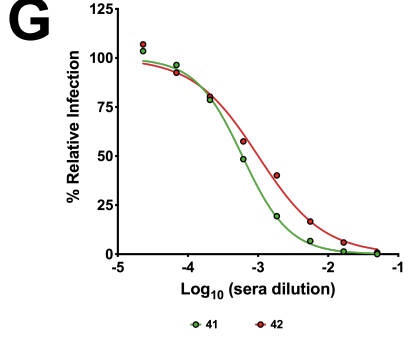
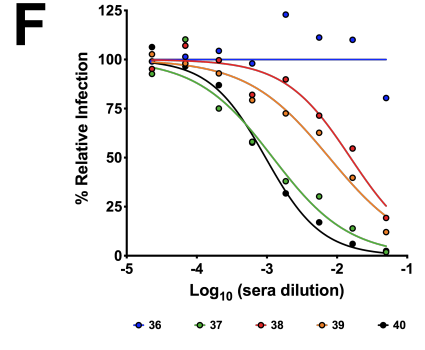
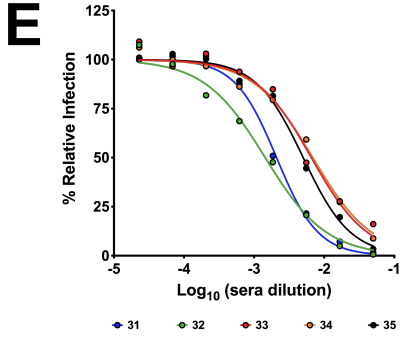
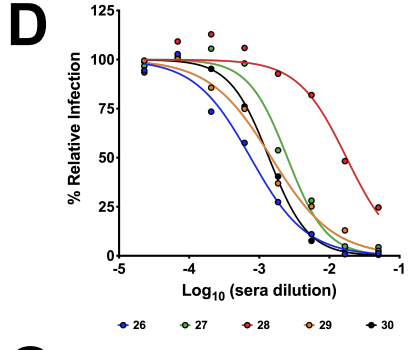
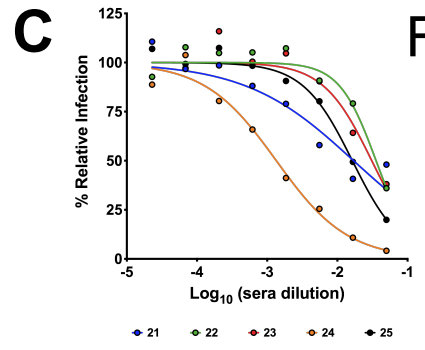
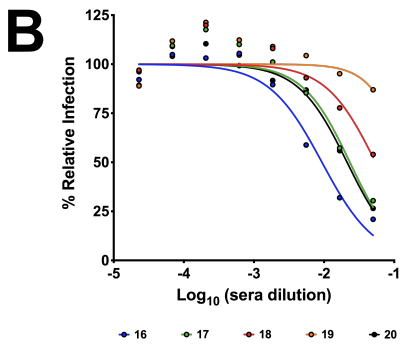
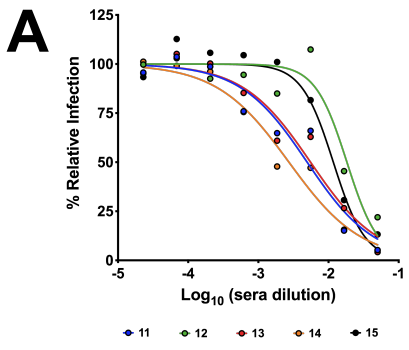
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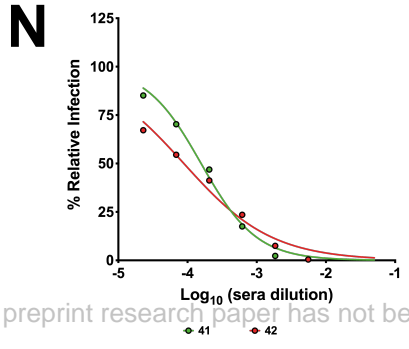
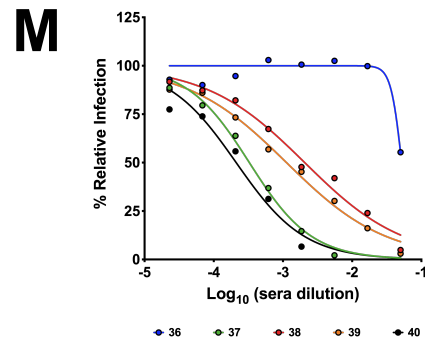
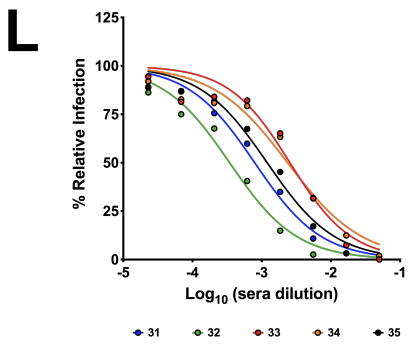
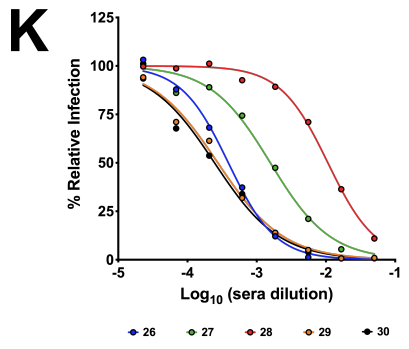


- VSV-SARS-CoV-2-S_{Δ21} + hACE2-Fc
- ▲ VSV + hACE2-Fc
- VSV-SARS-CoV-2-S_{Δ21} + mACE2-Fc
- ◆ VSV + mACE2-Fc

SARS-CoV-2



VSV-SARS-CoV-2-S_{Δ21}



Supplemental Data.

Antigenomic (+ sense) sequence of VSV-eGFP-SARS-CoV-2-S_{Δ21}:

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